

Department Mol. Biol
Subject 615/95 - 9/14/95
Name ANN KIM # 10
Address _____

Nationel® Brand

Computation Notebook

11 3/4" x 9 1/4", 4 x 4 Quad., 75 Sheets

43-648



0 73333 43648 8



AVERY
DENNISON

Office Products
Chicago, IL 60622

Ruben EXHIBIT #93

Department MOL. BIOL.
Subject 615195 - 9/14/95
Name ANN KIM # 10
Address _____

Hatfield® Brand

Computation Notebook

11 1/4" x 9 1/4", 4 x 4 Quad., 75 Sheets

43-648



0 73333 43648 8

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Ruben EXHIBIT 2093
Ruben v. Wiley et al.
Interference No. 105,077
RX 2093

27

6/23/95

1) Isolate 100ml 1Bt Amp Kan
 3ml A7 uninduced
 Incubate 37°C until OD₆₀₀ = 0.4-0.6
 2 1/2 hours w/ aeration
 Add 100mM IPTG to 2mM
 2ml
 Incubate 37°C w/ aeration 4 hours
 Spin culture 15 min 3K
 Resuspend pellet 10ml 1ell Gm HCl pH 8
 Store 4°C over weekend

6/26/95

Isolate HTPANX8804 5tbp ATG + PDI
 in 6M Gm HCl pH 5 in Acrylamide
 preparative gel

in 450ul H₂O
 Add 50ul Protein
 50ul 0.15% Na-DK
 75ul 50% TCA

Mix well

Spin 10 min

Resuspend pellet in 25ul 0.2M NaOH

Combine into 2 Tubes

Add equal volume of 2X Disoclation

Buffer

Heat 100°C 5 min

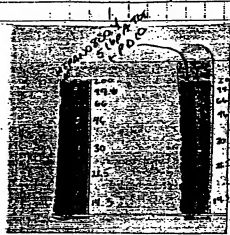
Run 100 V 1 1/2 hours

Stain + destain marker + part of gel

Cut desired band from gel

Cut up gel slice - ready to ship

6/26/95

pg 141
Product 200
PMS 29

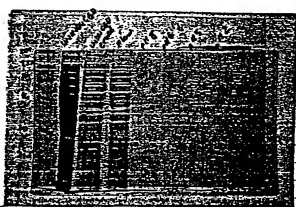
Send for AB
production
to Pecos farm
Cindy Haak
~ 0.75mg Abundance
~ 0.5mg eluted from
Immunodig

Spin Crude extract of HMSAF22A2
8K 20 min
Transfer supernatant to fresh tube
Prepare NH_4SO_4 Column.
2ml Buffer to make 1ml bed.
Wash Column 10ml H_2O
Equilibrate with 20ml 0.1M GnHCl pH 8
Add supernatant. Collect Flow
Wash column 30ml 0.1M GnHCl pH 8
Collect pH 8
Wash Column 30ml 0.1M GnHCl pH 6
Collect pH 6
Elute Protein 5ml 0.1M GnHCl pH 5
Collect eluted
Strip Column 30ml 0.1M GnHCl pH 2
Collect pH 2
Prepare to run on 12% Stacking
PAGE Gel

6/26/79

450 μ l H₂O
 50 μ l Protein
 50 μ l 0.15% NaDOC
 75 μ l 50% TCA

Mix well
 Spin Down
 Remove Supernatant
 Resuspend pellet 240 μ l 0.2N NaOH
 Add 20 μ l 2x Discontinuity Buffer
 Heat 100°C 5 min
 Run 20 μ l on gel w/Na MW Marker
 150V 1 1/2 hrs
 STAIN / DESTAIN



Does not look
 like anything
 induced

Remake most
 fragment

PR Fragment
 11749 + 11848
 10x #2
 10x BSA
 H₂O
 Xba I

20 μ l
 6
 6
 27
 60

incubate 37°C
 overnight

6/27/95

Inoculate 5ml LB+ Amp+Kan
 with frozen glycerol stock
 HTPA 198 5/10 + 6 PD10 D5
 incubate 37°C w/ aeration 2 1/2 hours
 Inoculate 5ml LB+ Amp/Kan w/ culture
 add 100ul 100mM IPTG to culture
 incubate all 37°C w/ aeration
 Inoculate 50 ml LB+ Kan/Kan
 to do large scale induction

PCR Fragment
 Kba Digest

PPT with Ethanol and NaOH
 Spin 10 min
 Remove supernatant
 Wash pellet 70% Ethanol
 Spin 5 min
 Remove supernatant
 Allow pellet to dry
 Resuspend pellet 44ul TE
 Add 5ul Buffer
 1ul Sp6
 let digest 37°C 6 hr

6/28/95

Inoculate 100 ml LB+ Amp/Kan
 with 35 ml overnight culture
 of HTPA 198 504 Strp ATG + PD10
 D5
 OD₆₀₀ ~ 0.87
 incubate 37°C w/ aeration for 2 hrs
 until OD₆₀₀ ~ 0.4-0.6

6/28/95

Run.

- 3x [
- 1 - Rainbow Marker
 - 2 - HTPAN08S04 51bp ATG + PD10 UNINDUCED
 - 3 - " " " " " INDUCED
 - 4 - HBSAF22 + PDE UNINDUCED
 - 5 - HTPAN08S04 51bp ATG - pH5.

Run 150 V: 1 1/4 hours.

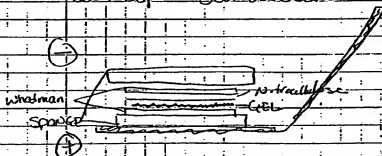
Set up transfer.

Transfer Buffer:

25 mM Tris pH 8.3
 192 mM Glycine
 20% METHANOL.

4°C

Set up: Sandwich.



Run 100 mV
 40 min
 Run 200 mV
 20 min

Store blots in Blocking Buffer
 WESTERN BLOCKING BUFFER:

3% BSA Fraction V
 0.02% Na AZIDE
 in PBS

at 4°C

34
 M

10/29/95

Run 2nd of Fragment with 1 Kb ladder



fragment looks good

Set up ligations at 21°C

3 Nov/95 S.H.I.

	1	2	3	4	5	6
DNA Fragment	2	2	2	1	1	1
PGE30 Nco/SphI	1	1	1	1	1	1
PGE30 SphI/Nco	1	1	1	1	1	1
10X Buffer	2	2	2	2	2	2
T4 Ligase	1	1	1	1	1	1
H ₂ O	14	14	15	16	16	17

Incubate reactions 16°C overnight



HTPA208504 81bp ATG + PD10 DS protein

Spin 8K 30 min

Transfer Supernatant to fresh tube.

Crude Extract

Prep. W. Sep. Column

6/2/92

Prepare Bed - 3ml
 Strip Column - 30ml 0.2N NaOH
 Wash 50ml H₂O
 Charge 50ml 0.1M NaSO₄
 Wash 50ml H₂O - Equilibrate 50ml 0.1M NaOH
 Add Supernatant to column
 Collect - flow
 Wash 40ml 0.1M HCl pH 2
 Collect - pH 8
 Wash 40ml 0.1M HCl pH 6
 Collect - pH 4
 Elute 10ml 0.1M HCl pH 5
 Collect - pH 5
 Strip - 40ml 0.1M HCl pH 2
 Collect - pH 2

Run fraction on Gel

450ul H₂O
 50ul Fraction
 50ul 0.15% NaDOC
 75ul 50% TCA

Mix well

Spin 10 min

Remove Super natant

Resuspend pellet with 0.2N NaOH

Add 10ul 2X Discontinuation Buffer

Heat 100°C 5min

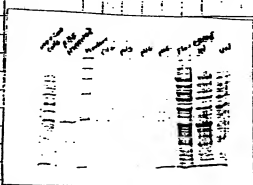
Spin 5min

Run all on 12% Stackin

PAGE Gel 150V 1hr

Stain 30min / DESTAIN 30min

6/30/95



looks like most of
the plasmid did
not stick into
column -
try Reapplying
crude extract to
Flav. box
aka culture finally
changed again

Transform ligations

Thaw Chemically Competent M15 cells
on ice

To 100 μ l thawed cells add 10 μ l of
ligation

1st set on ice 1 hour

Heat 42°C 45 sec

Put on ice 2 min

add 400 μ l LB to tube

Incubate 37°C 1 hour

plate 100 μ l into LB + Amp/Kan
plates

incubate at room temp over
weekend

7/3/95

Received primers for HTPAN08504
into pCDNA 3' HA TAG
5' HA TAG
→ Naitang

#12305 HTPAN08504
5' Bam HI + 3' 5' Untranslated Region
for 3' HA TAG in pCDNA.

GCG GGC GGA TCC TGC CTG GCT GAG CTTA CAG
CAATC

#12306 HTPAN08504
3' Xba I for 3' HA TAG in pCDNA
No Stop Codon

GCG GGC CCA GAG GCG GGA AAC TAA AAA GGC
CCC GAA AAA ACT G

#12307 HTPAN08504
5' Bam HI for 5' HA TAG and Regular pCDNA

GCG GGC GGA TCC GGT ATG ATG GAG GTC CAG GGG
GTA C

#12308 HTPAN08504
3' Xba I for 5' HA TAG and Regular pCDNA
has restriction site

GCG GGC CCA GAG TTA TTA GCG AAC TAA AAA
GCG CC GAA AAA AC

PCR products

07/13/85

Set-up PCR

3' HA TAG

12805	1
12806	6
10x dNTP	10
10x PCL	10
Taq	0.4
H ₂ O	60.4
DNA	1
	100.2

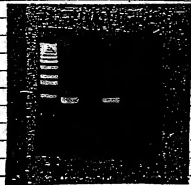
5' ATTAG

10x		10x
60	12807	37
60	12808	6
100	10x dNTP	10
100	10x PCL	10
	Taq	0.4
	H ₂ O	68.9
	DNA	1
		100.2

PCR

95°C 5 min
 95°C 30 sec
 65°C 30 sec } 25x
 72°C 1 min
 72°C 9 1/2 min
 4°C Hold

Run T7 on gel



Looks good
 Combine Tubes

Dilute plate with equal volume PEG/salt
 Spin 10 min
 Remove & supernatant
 Wash pellet 1000 µl 70% Ethanol
 Spin 5 min
 Remove supernatant
 Wash pellet 10 min
 Dry at RT

Resuspend pellet in a total of
100 μ l TE
Remainder left on gel



Store -20°C Fragment #3
Box

(10 μ l)

Pick colonies from transformation
(4/30) into LB + Amp/Kan - HMSA22

- 46 ϕ #1 - 2 ϕ #3
- 46 ϕ #2 - 2 ϕ #5

Inoculate 57 μ l w/acerol: 4 hours
Set up PCR

		10X
5' PDE Vector	0.2	20
3' PDE Vector	0.14	14
10X dNTP	3.2	320
10X PCR	3.2	320
H ₂ O	23.7	2310
Taq	0.16	16
Culture	2	
	32.0	3000 - 30 μ l / tube

		100X
5' SpHMSA22	1.6	160
3' PDE Vector	0.14	14
10X dNTP	3.2	320
10X PCR	3.2	320
H ₂ O	21.7	2170
Taq	0.16	16
Cult.	2	
	32.0	3000 30 μ l / tube

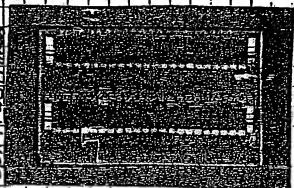
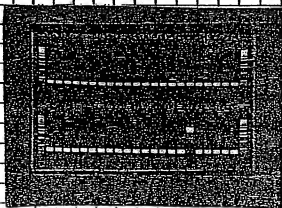
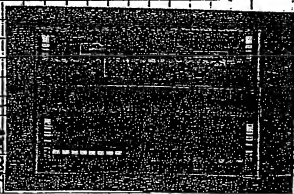
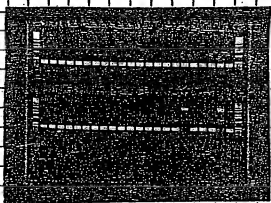
40

7/3/95

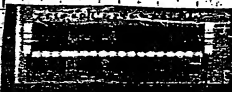
PCR Program #66

95°C	5 min	} 30x
95°C	30 sec	
72°C	20 sec	
72°C	1 min	
72°C	7 1/2 min	
4°C	Hold	

7/4/95 Holiday

7/5/95 Run 10 ul of PCR reaction
with 1Kb ladder - HIN3AF 22

Set up PCR 11/1/95



Does not look like
the cloning worker.
Try Redtransforming
w/ 7/95

Set up Dyeing in 6 HTPM 08504
for pc-DNA (p37A)

5' HA TAG

Fragment	20
10x #2	5
Prim HT	1
Kho T	
H ₂ O	23
	<u>50</u>

3' HA TAG

Fragment	20
10x #5	5
Prim HT	1
Vha T	
H ₂ O	23
	<u>50</u>

Incubate 37°C overnight

7/9/95

Pick more colonies from 6/9/95 - HMTAF 22
into LB + Amp
Incubate 37°C 4 hours

Set up PCR

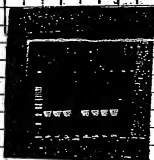
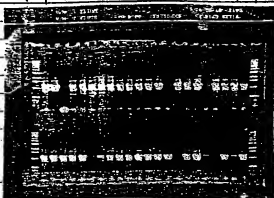
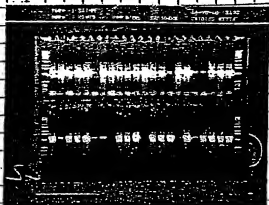
5' Sense	1.1	160
3' PDB	0.11	14
10x HNTA	3.2	320
10x PCR	3.2	320
Taq	0.16	16
H ₂ O	826.617	2170

07/16/95

PDR Proxy 66

HMSAP22

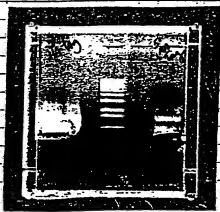
Run 10 million GEL w/ 1 Kb ladder



looks good
 incubate 120 min @
 1.8 x 10⁶ / ml
 with (P) and 1.8 x 10⁶ / ml
 incubate 30 min @
 37°C w/
 aeration in air

1 MP Gel Purification
 HPLC 3' HA 1.8
 5' HA 1.8

Load onto 0.8% EMP Gel
 Cut out all slices
 for Taka Pfu
 Near clean fragment



HCPAN 2501

H/G/95

Gene Clean

Add 1000 μ l AbT

Heat 55°C 5min

Add 10 μ l Glass milkLet incubate at RT
5min

Spin 10 sec

Remove Supernatant

Resuspend pellet in

SDSul Wash Buffer

3X Spin 10 sec

Remove Supernatant

Spin 10 sec

Remove Supernatant

Resuspend pellet 40 μ l TE

Heat 55°C 5min

Spin 1 min

Transfer Supernatant to next tube

Add Resuspend pellet 20 μ l TE

Heat 55°C 5min

Spin 1 min

Transfer Supernatant

Run 1 μ l of Fragment on gel with1 kb ladder
Store -20°C in Fragment #5 Box

Will need Vector to do ligations
None present in correct
Digestions. So set up Digests

CDNA 5' HA	10 μ l
10x2	5
H ₂ O	23
Prim 1/2 I	1/1
	50

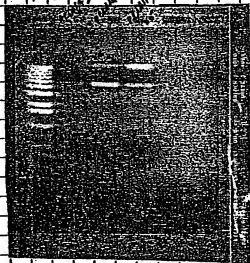
CDNA 3' HA 6'	17
10x1	5
H ₂ O	26.0
Prim 1/2	1/1
	50

7/6/95 pCDNA-I 1.1
 10X #2 5
 H₂O 41.9
 Barn 1A1 11
 50

incubate 37°C overnight

7/13/95

Remainder of 196 Digest on Gel



pCDNA Should be
 24.7 Kb

pCDNA 5'45' HA
~~fragments~~
 plus DNaase. insert
 1600 bp

pCDNA 3 fragments

looks like not completely
 digested.
 Add 1/2 more. Enzyme
 incubate 37°C. well
 on top of 37°C over
 weekend.

Rec'd Clone from Sigis

Clone I.D. HE2PM 2.1 #2 (RB)

Requested By Ann Kim

Transfer Folder 07/05/95 + transfr-2

Processed By Luis Rodriguez

1/1/95

7/7/96

Small scale inductions
 - from overnight cultures from p42.
 Inoculate 200 μ l LB + Amp Kan. w/ 15 μ l of Cult. 100 ml
 incubate 37°C w/ aeration 2 hrs.
 OD₆₀₀ = 0.4 \pm 0.05
 Add 100 ml IPTG to 200 ml - 4 ml
 incubate 37°C w/ aeration 4 hrs.
 Spin culture 10 min.
 Remove supernatant
 Resuspend pellet in 100 μ l H₂O
 Add 100 μ l of Dissociation Buffer
 Heat 100°C 5 min
 Spin 2 min
 Run 5 μ l on gel with marker and
 uninduced cultures as 15 well
 on 12 to 200 base pairs gels.
 Run 150 V 1.2 amps
 Stain overnight at RT.

7/10/96

Run PCR/ Digest on 0.8% LMP gel
 with 16 lanes

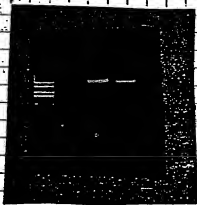


Get out Gel. Slides
 Clean. Clean

Resuspend Gel. Slides
 200 μ l Dist. H₂O
 Heat 55°C 5 min
 Add 3 μ l glass beads
 hit pot. at RT 2-3 min
 w/ occasional mixing
 Spin 10 sec

7/10/95

Remove Supernatant
 Resuspend pellet 100 μ l Wash Buffer
 Spin 10 Sec
 Remove Supernatant
 2x { Add 100 μ l Wash Buffer
 Spin 10 Sec
 Remove Supernatant
 Spin 10 Sec
 Remove Supernatant
 Resuspend pellet 15 μ l TE
 Heat $+55^{\circ}\text{C}$ 1 min
 Spin 10 Sec
 Transfer Supernatant to fresh tube
 Resuspend pellet 15 μ l TE
 Heat $+50^{\circ}\text{C}$ 1 min
 Spin 10 Sec
 Transfer Supernatant to New tube
 Run 2 μ l on gel with 1 μ l ϕ ladder



Vectors look good
 GATE

store -20°C
 Vector #2 BR

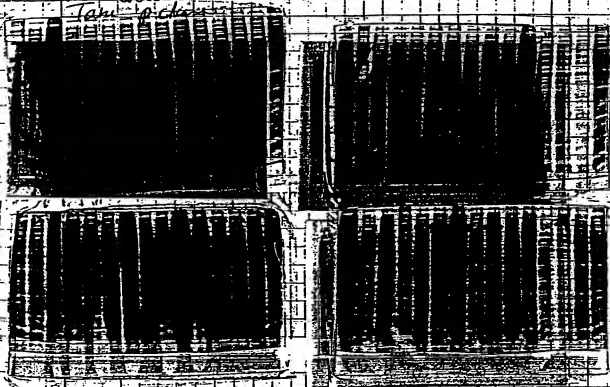
Set up ligations
 10 μ l HEPAN 0.8504
 5' + 3' HA Tags
 Phosphite
 (pg 18)

7/10/95

	1	2	3	4	5	6	7	8	9
HTPAND504	1								
5' HA-tag Bam/Kho	1	1	1	1					
HTCPA106504									
3' HA-tag Bam/Kho			1		1				
PCDNA Bam/Kho	4					4			
PCDNA 5' HA Bam/Kho		3					3		
PCDNA Bam/Kho 3' HA			2	1				2	
10x Buffer	2	2	2	2	2	2	2	2	2
1x O.D.	12	12	12	16	16	13	14	15	17
1x Ligase	1	1	1	1	1	1	1	1	1

Incubate 16°C overnight

DESTAIN Gels 33°C - 2 hrs. - HMSA P22 P8670



7/11/95

	5' HA	3' HA	pcDNA
DAADNA	17	27.8	11.8
10X BAW	5	5	5
10X BSA	5	5	5
H ₂ O	22	11.2	37.2
Enzyme	11	1	1
	50	50	50.1

Digest with Xba I and Bam HI

Incubate 37°C O/N

HTP AND SO4 pcDNA constructs

ligation from 7/10/95 pg 47

Transform ligations (100 µl) into
100 µl of Chemically Competent
DH5α cells.

Thaw Cells on ice

Aliquot 100 µl into Sterile tubes

Add 10 µl of ligations

Mix by pipetting

Incubate on ice 1 hour

Heat 45°C 45 sec

Place on ice 2 min

Add 400 µl LB

Incubate 37°C 1 hour

Plate 100 µl + 200 µl into

LB Amp plates

Incubate 37°C overnight

7/12/95

Pick 4 transformed cells into 1 RT Amp
 HTPANORSOY + pCDAM constructs

- 1 HTPANORSOY + pCDMA
- 2 HTPANORSOY + 5' HA Tag pCDNA
- 3 HTPANORSOY + 3' HA Tag pCDNA

Incubate 30°C w/ 1000 rpm 4 hrs
 Set up PCR

for #1 & 2

for #3

H12307	2
FP13	0.01
1000 RPM	3.2
10x PCR	13.2
Taq	0.15
H ₂ O	21.4
Culture	2
	32

H2305	2.5
FP13	0.01
1000 RPM	3.2
10x PCR	3.2
Taq	0.15
H ₂ O	20.9
Culture	2
	32

PCR Prog 6L

① HTPANORSOY
 Plasmid
 ② H₂O only

95°C	5 min	} 30s
95°C	20 sec	
55°C	20 sec	
72°C	1 min	
72°C	7 1/2 min	
4°C	Hold	

Run 10ul on Gel with 1 kb ladder

No positives seen - Not even (+)
 plasmid PCR'd. Rado tomorrow
 with other primers

Casare Out 07/13-07/17

2/13/95

Add Equal Volume PEG / NaCl
4% 0.1M

PCR

HTPANO8 - PC DNA C-mutants

102

3

12307

2

12305

3

12308

3

12306

3

10x dNTP

3.2

10x dNTP

3.2

10x PCR

3.2

10x PCR

3.2

H₂O

18.4

H₂O

18.4

Taq

0.2

Taq

0.2

Cult. vol.

2

Cult. vol.

2

3.2

3.2

3.2

3.2

Rem PCR

Prog

Col

5' Kp

Kp

Kp

Galacto Kinase Assays

DH₅ Galactose + ATP $\xrightarrow{\text{GalK}}$ α -D-Galactose-1-phosphate + ADPADP + PEP $\xrightarrow{\text{PepK}}$ Pyruvate + ATPPyruvate + β -NADH $\xrightarrow{\text{LacK}}$ Lactate + β -NADOD₃₄₀

Reagents:

A:

100 mM Potassium Phosphate Buffer pH 7.0

B:

100 mM D-Galactose

C:

5.9 mM Adenosine 5 Triphosphate
(ATP)

Add 13% PEG 8000 / 1.6M NaCl = 850 ml
 mix well.
 Store -20 °C O/N.

7/13/95

Ratio of Galactose

λ 800	30-180 sec	λ 800	30-180 sec	λ 800	30-180 sec
H	$0.682 - 0.681$ 150 sec	1-2	$0.6919 - 0.6792$ 150 sec	1-3	$0.7327 - 0.734$ 40 sec
2-1	$-6 \times 10^{-4} / \text{sec}$ $0.6665 - 0.5709$ 150 sec	2-2	$1.1 \times 10^{-4} / \text{sec}$ $0.6604 - 0.6409$ 150 sec	2-3	$-3 \times 10^{-5} / \text{sec}$ $0.8018 - 0.798$ 40 sec
3-1	$0.4 \times 10^{-4} / \text{sec}$ $0.706 - 0.703$ 80 sec	3-2	$9.3 \times 10^{-4} / \text{sec}$ $0.6777 - 0.6701$ 150 sec	3-3	$2.8 \times 10^{-4} / \text{sec}$ $0.7750 - 0.7730$ 40 sec
4-1	$3.5 \times 10^{-5} / \text{sec}$ $0.6335 - 0.5931$ 150 sec	4-2	$1.3 \times 10^{-4} / \text{sec}$ $0.6203 - 0.6192$ 150 sec	4-3	$-5 \times 10^{-9} / \text{sec}$ $0.7554 - 0.7587$ 40 sec
5-1	$6 \times 10^{-4} / \text{sec}$ $0.6541 - 0.5678$ 150 sec	5-2	$9.4 \times 10^{-4} / \text{sec}$ $0.6273 - 0.6196$ 150 sec	5-3	$3 \times 10^{-4} / \text{sec}$ $0.7937 - 0.7885$ 40 sec
Depth Blank	$7.8 \times 10^{-4} / \text{sec}$ $0.6105 - 0.1350$ 150 sec	Blank	$9.0 \times 10^{-4} / \text{sec}$ $0.7280 - 0.7212$ 150 sec	Depth Blank	$5.8 \times 10^{-5} / \text{sec}$ $0.533 - 0.1645$ 40 sec
	$3.2 \times 10^{-3} / \text{sec}$		$4.15 \times 10^{-5} / \text{sec}$		$4 \times 10^{-3} / \text{sec}$

7/14/95

Pi-PER HYDROLYSIS pcDNA constructs
 Run on gel with 1 Kb ladder

See pg 54

7/13/95

H5/H9	0.7705 - 0.6006	H9 DT	0.3019 - 0.6740
DT - 20°	150 sec	E5 - 20°	150 sec
7/14/95	1.8×10^3 / sec	7/14/95	1.85×10^3 / sec
Galk + 10%	0.6771 - 0.1012	Blank	0.8696 - 0.0053
Galk - 20°	150 sec		150 sec
7/13/95	3.8×10^3 / sec		2.9×10^3 / sec

Inoculate 5ml TB + Amp
with HTPANX8 pcDNA Cultures

- 1: A1, A2, A3, A4
- 2: D1, D2, D3, D4, E1, E2, E5
- 3: E9, E10, E2, G1, G2, H3, H4

Inoculate 37°C w/ aeration 9/1N

7/18/95

Baking Mann pups - HTPANX8 pcDNA construct

Spin 1ml Cultures

Remove Supernatant

Resuspend pellet 800ul STE7 + RNaseT

lysozyme

Heat 1 sample 70°C 1 min

Spin 5 min

Resuspend pellet

Add 800ul 15% PEG 800 / 1.0M NaCl - mix well

Spin 10 min

Remove Supernatant

Wash pellet 1 ml 70% Etanol

Spin 5 min

9/8/95

Remove Supernatant
 All this pulled to bag at RT 10 min
 Resuspension pulled in 250 μ l TE
 Rinse bulb and gel with 1 kb
 Add

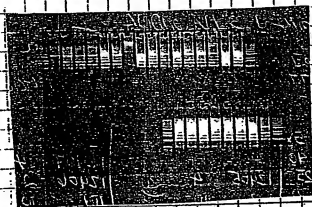


Setup Digestion:
 XhoI 10 min

		10x
DNA	10	
10xHF2	3	50
H ₂ O	10.8	302.4
Boam	0.1	1.8
Xho	0.8	1.8
	30	

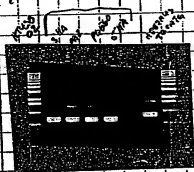
Incubate Digests at 37°C O/N

Rinse HUSAF22 + PDE Digests on 10% Gel
 with 1 kb ladder - Band II / Sph I



Cannot tell
 anything with
 this digest
 - Too much Chromosomal
 - Try Digests with
 EcoRI and HinfI

4/19/95

Run 1/1 of fragment on gel with
11 kb ladder

HG55H43 17+ATG
 ready for 11/1
 store 20 Fragment
 #2 Box

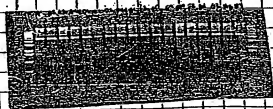
H43

H43B02 Fragments
Set up Digests

3HA 12408+12409				R5HA 12406+12407				PA7 12404+12405				P0E 12402+12403			
DNA	10	DNA	10	DNA	10	DNA	10	DNA	10	DNA	10	DNA	10	DNA	10
10x	5	10x	5	10x	5	10x	5	10x	5	10x	5	10x	5	10x	5
H2O	34	H2O	34	H2O	34	H2O	34	H2O	34	H2O	34	H2O	34	H2O	34
Boon		Yho	1	Boon	1	Yho	1	Boon	1	Yho	1	Boon	1	Yho	1
	50		50		50		50		50		50		50		50

Incubate 37°C 4 hrs - Add 10xTaq
 enzyme and incubate ON at 37°C

H43B02 - PCR DNA Constructs - form/clone digests



7/20/95

HTRADIX Mini pups

Spin 5ml culture

8K 15 min

Resuspend pellet in Resuspension

Buffer + 10ml

Add 10ml Lysol B. 60

mix gently

let sit 15 min at RT

Add 10ml Neutralization Buffer

mix well

let sit on ice 20 min

Spin 8K 30 min

Transfer supernatant to flat tube

Add 10ml 30% PEG and 10ml

5M NaCl

mix well

store 4°C 20 min

Spin 8K 30 min

Pour 50 Supernatant

Resuspend pellet 10ml 70% Ethanol

Spin 8K 15 min

Pour 50 Supernatant

Add 10ml pellet to 2 x 400ml TE

Resuspend in 2 x 400ml TE

precipitate DNA

100% Vol 3M Na acetate 2x Vol 100% Ethanol

store -20°C O/N

No 1/20/95

7/21/95

Pak Colonies into 1B+ Amp Media

1-24

5-12

9-48

2-24

10-12

10-96

3-12

7-24

4-12

8-24

75

7/24/95

Clones look good -
Monday..... inoculate for miniprep

HTRANOS

Spin DNA 10min

Pour off Supernatant

Wash pellet 2x 70% Ethanol

Spin 5min

Pour off Supernatant

Put pellet Dry at RT

Resuspend in 100ul of TE Total

Ready to clean up

9/24/95

Inoculate 5ml TB + Amp w/
Cultures from HTRANOS 1.0g

1- HTRANOS

3' HATAG B/X

- A1, A2, A3, A4

2

3' HATAG B/X/B

- C1, C2, C3, C4

3

5' HATAG B/X

- E1, E2, E3, E4

4

5' HA Tag X/B

- F5, F6, F7, F8

5

pCDNA B/X

- G1, G2, G3, G4

6

pCDNA X/B

- H5, H6, H7, H8

7

PA2 B/X

- A1, A2, A3, A4

8

PA2 X/B

- C5, C6, C7, C8

9

PA2 X/B

- C9, C10, C11, C12

10

PA2 X/B

- C13, C14, C15, C16

11

PA2 X/B

- C17, C18, C19, C20

12

PA2 X/B

- C21, C22, C23, C24

13

PA2 X/B

- C25, C26, C27, C28

14

PA2 X/B

- C29, C30, C31, C32

15

PA2 X/B

- C33, C34, C35, C36

16

PA2 X/B

- C37, C38, C39, C40

17

PA2 X/B

- C41, C42, C43, C44

18

PA2 X/B

- C45, C46, C47, C48

19

PA2 X/B

- C49, C50, C51, C52

20

PA2 X/B

- C53, C54, C55, C56

21

PA2 X/B

- C57, C58, C59, C60

22

PA2 X/B

- C61, C62, C63, C64

23

PA2 X/B

- C65, C66, C67, C68

24

PA2 X/B

- C69, C70, C71, C72

25

PA2 X/B

- C73, C74, C75, C76

26

PA2 X/B

- C77, C78, C79, C80

27

PA2 X/B

- C81, C82, C83, C84

28

PA2 X/B

- C85, C86, C87, C88

29

PA2 X/B

- C89, C90, C91, C92

30

PA2 X/B

- C93, C94, C95, C96

31

PA2 X/B

- C97, C98, C99, C100

32

PA2 X/B

- C101, C102, C103, C104

33

PA2 X/B

- C105, C106, C107, C108

34

PA2 X/B

- C109, C110, C111, C112

35

PA2 X/B

- C113, C114, C115, C116

36

PA2 X/B

- C117, C118, C119, C120

37

PA2 X/B

- C121, C122, C123, C124

38

PA2 X/B

- C125, C126, C127, C128

39

PA2 X/B

- C129, C130, C131, C132

40

PA2 X/B

- C133, C134, C135, C136

41

PA2 X/B

- C137, C138, C139, C140

42

PA2 X/B

- C141, C142, C143, C144

43

PA2 X/B

- C145, C146, C147, C148

44

PA2 X/B

- C149, C150, C151, C152

45

PA2 X/B

- C153, C154, C155, C156

46

PA2 X/B

- C157, C158, C159, C160

47

PA2 X/B

- C161, C162, C163, C164

48

PA2 X/B

- C165, C166, C167, C168

49

PA2 X/B

- C169, C170, C171, C172

50

PA2 X/B

- C173, C174, C175, C176

51

PA2 X/B

- C177, C178, C179, C180

52

PA2 X/B

- C181, C182, C183, C184

53

PA2 X/B

- C185, C186, C187, C188

54

PA2 X/B

- C189, C190, C191, C192

55

PA2 X/B

- C193, C194, C195, C196

56

PA2 X/B

- C197, C198, C199, C200

57

PA2 X/B

- C201, C202, C203, C204

58

PA2 X/B

- C205, C206, C207, C208

59

PA2 X/B

- C209, C210, C211, C212

60

PA2 X/B

- C213, C214, C215, C216

61

PA2 X/B

- C217, C218, C219, C220

62

PA2 X/B

- C221, C222, C223, C224

63

PA2 X/B

- C225, C226, C227, C228

64

PA2 X/B

- C229, C230, C231, C232

65

PA2 X/B

- C233, C234, C235, C236

66

PA2 X/B

- C237, C238, C239, C240

67

PA2 X/B

- C241, C242, C243, C244

68

PA2 X/B

- C245, C246, C247, C248

69

PA2 X/B

- C249, C250, C251, C252

70

PA2 X/B

- C253, C254, C255, C256

71

PA2 X/B

- C257, C258, C259, C260

72

PA2 X/B

- C261, C262, C263, C264

73

PA2 X/B

- C265, C266, C267, C268

74

PA2 X/B

- C269, C270, C271, C272

75

PA2 X/B

- C273, C274, C275, C276

76

PA2 X/B

- C277, C278, C279, C280

77

PA2 X/B

- C281, C282, C283, C284

78

PA2 X/B

- C285, C286, C287, C288

79

PA2 X/B

- C289, C290, C291, C292

80

PA2 X/B

- C293, C294, C295, C296

81

PA2 X/B

- C297, C298, C299, C300

82

PA2 X/B

- C301, C302, C303, C304

83

PA2 X/B

- C305, C306, C307, C308

84

PA2 X/B

- C309, C310, C311, C312

85

PA2 X/B

- C313, C314, C315, C316

86

PA2 X/B

- C317, C318, C319, C320

87

PA2 X/B

- C321, C322, C323, C324

88

PA2 X/B

- C325, C326, C327, C328

89

PA2 X/B

- C329, C330, C331, C332

90

PA2 X/B

- C333, C334, C335, C336

91

PA2 X/B

- C337, C338, C339, C340

92

PA2 X/B

- C341, C342, C343, C344

93

PA2 X/B

- C345, C346, C347, C348

94

PA2 X/B

- C349, C350, C351, C352

95

PA2 X/B

- C353, C354, C355, C356

96

PA2 X/B

- C357, C358, C359, C360

97

PA2 X/B

- C361, C362, C363, C364

98

PA2 X/B

- C365, C366, C367, C368

99

PA2 X/B

- C369, C370, C371, C372

100

PA2 X/B

- C373, C374, C375, C376

101

7/24/95

Transform ligations from 7/20/95
 ligations P_{H3V} - 18 - H74802 + PCE60

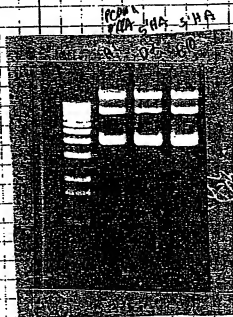
Throw M15 up 4 cells on ice
 Add 100ul of Chemically Competent
 cells to 10ul of ligation
 incubate on ice 0.5 hour
 heat 42°C 45 sec
 place on ice
 add 400ul LB
 incubate 30 min 1 hour
 plate 100 + 250 ul onto LB Amp/Km.
 plates
 incubate at 37°C O/N.

HTPABOS pDNA Constructs

2x Phenol/Seaq (1.0M extract)
 5x Seaq extract
 PPT DNA
 710 ul 3M Na acetate pH 5.3
 2.1 ul 100% ethanol
 Spin 10 min
 Pour off supernatant
 Wash pellet 1000ul 70% ethanol
 Spin 5 min
 Pour off supernatant
 Allow pellet to dry at RT 10
 min
 Resuspend pellet 200ul TE
 Run 1ul on gel with 100 ladder

77

7/24/95



Plasmid looks good.
Set up Digest

DNA	4
10X #2	3
H ₂ O	22
Bam HI	0.5
Xho I	0.5
	<hr/> 30

Incubate 37°C O/N.
Run on gel tomorrow.

7/25/95

RADIATION PROBLEM FOUND -
- UNABLE to travel freely around
3rd floor 9670

- Take plate out at 37°C
H₂O + 15002 + 10000

- Take culture tubes at 37°C
H₂O + 15002 + 10002 + 10002

- Boiling mixes

Spin 2ml culture 5 min

Remove supernatant

Resuspend pellet 1000 µl STEA

Resuspend pellet 1000 µl STEA

Heat 100°C 2 min

Spin 10 min

7/25/95

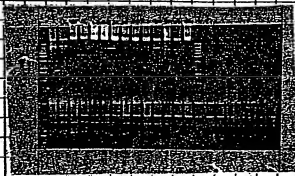
Remove Pellet
Add equal volume 18% PEG/NaCl
mix well
Stir - 20°C

7/26/95

NO WORK + Cleaning of
of Radioactivity by RSO

7/27/95

Spin B. Mon. S. 15 min
Remove supernatant
Wash pellet 1.5 ml 50% ethanol
Spin 5 min
Remove supernatant
Allow Pellet to dry at RT 15 min
Resuspend in 200 µl TE
Run 2nd on gel with 1 kb ladder.



1-8 HT4502 + 3'HA CAN
H1, A2, A3, A4, C1, C2, C3, C4
9-16 HT4502 + 5'HA CAN
E1, E2, E3, E4, F1, F2, F3, F4
17-24 HT4502 + 3'HA CAN
G1, G2, G3, G4, H1, H2, H3, H4
25-32 HT4502 + PAZ
A1, A2, A3, A4, C1, C2, C3, C4
33-36 HT4502 + PAZ
37-40 HT4502 + PAZ
41-44 HT4502 + PAZ
45-48 HT4502 + PAZ
49-52 HT4502 + PAZ
53-56 HT4502 + PAZ

7/27/95

Set up Digests

1, 24 + 33, 34, 35

Digest Bam/Xba

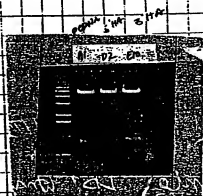
25 - 32 + 36

Digest with Bam/Xba

DNA	10	29X	DNA	10	10X
10X #2	3	87	10X #2	3	30
H ₂ O	16.8	487.2	Bam H1	0.1	1
Bam H1	0.1	2.9	Xba I	0.1	1
Xba I	0.1	2.9	H ₂ O	16.8	168
30ul		20ul tube	30ul		24.1/ul

Incubate at 27 - 3 hrs

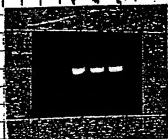
Run HTPAW Digests on gel w/ 1Kb ladder



Digests look good
 Submit for seq
 with internal
 primers

7/11/95

Run ~~1000~~ of plasmid on gel w/ 1 kb
 plasmid looks good



Selling Digest

Bam / Xba I

DNA	1
H ₂ O	3
H ₂ O	25.6
BamHI	0.2
Xba I	0.2
	130ul

Incubate 37°C 6/11

Transfection

Remove Spent media
 Wash Cells in 3ml DMEM without
 Serum

make Transfection Cocktail

300ul	
30ul	Nutrient
0.75ul	DEAE Dextran (Zongshel Stock)
0.3ul	Chloroquine (Gibco final Stock)
25.8ul	DMEM

To: 4ml of cells Transfect in Cocktail
 Add 10ul DNA

HT10802	pcDNA	Constructs
HT10802	pcDNA	Constructs

8/30/95

Digest with Bam / Xba - HEZAM2.1 + PAZ

DNA	2
H ₂ O	15.6
10x ⁺ 2	2
BamHI	0.2
XbaI	0.2
	20ul

Incubate 37°C 2hrs
 Run 10ul on gel w/ 1kb ladder

HEZAM2.1 + PAZ
 Bam HI / Xba I

Submit for Sequencing with internal
 primers and with PAZ specific
 primers.

HTUSB02 + PAZ

Construct sent to Protein Expression
 Submit for Sequencing to Confirm

FAS Ligand - HTPALXX

Antibody - Rec of Rabbit Ab
 from Porcine farms

Ready for Western Detection

FAS LG	RABBIT	#11940	→ #1
		#11941	→ #2

Pre Bleed - 6/30/95
 Post Bleed #1 - 8/25/95

8/30/95

Western blots from 6/28/95
(See pg 313)Remove 24 blots from Blocking Buffer
To 10 ml Eyring blocking Buffer
Add 100 μ l ABC Solution 1:100
Incubate ON at RT Shaking

- #1 = Test bleed #1 - 11941
 #2 = Test bleed #1 - 11940
 3 = Prebleed #11940
 4 = Prebleed #11941

8/31/95

FAS LIG Western

Wash off
 Rinse 1st Ab
 Wash filters in 1x PBS
 Wash 10 min 10 ml 1x PBS 5 min
 Add at RT w/ shaking
 5th Ab - Rabbit Anti Rabbit
 Peroxidase - Dilute 1:2000 in 1x PBS
 Incubate at RT w/ shaking for
 1 hr
 Rinse filters 1x PBS
 Wash 1x in 10 ml 1x PBS - 5 min
 at RT w/ shaking
 Rinse filters 20 ml
 50 mM Na₂PO₄ pH 7 for
 5 min at RT w/ shaking

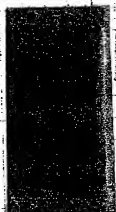
8/30/95

Add Substrate:

12.5 ml 50 mM $\text{Na}_2\text{H}_2\text{PO}_4$ pH 725.0 mg β -NADH25.0 ml Phenol Equilibrated
in TRIS pH 7.08.5 ml 30% H_2O_2 solution37.5 ml NBT - Nitro Blue Tetrazolium
(10 mg/ml)Incubate at RT till color change
develops.Stop Rxn with dH₂O.

Dry and Label blots

FAS Lig - WESTERN

Probed
#1941Probed
#1940

1° Ab.

Test Bleed #1
#1940Test Bleed
#1941

lane #1 - Rainbow Marker

#2 - HTPAN08504

51bp ATG in PD10

UNINDUCED

#3

HUSAF22

51bp ATG in PD10

INDUCED

#4

HUSAF22 + PQE70

INDUCED

#5

HTPAN08504

51bp ATG in PD10

purified

NTA Column - pH 5.0

See Pg 141

813195

PCR HTPANOS Constructs
POECO

12975	514p	2	12986	1854p	2
2845		0.4	2865		0.4
10A		10.1	10.1		10
10A		10	10.1		10
H2O		74.3	H2O		76.3
Tsp		0.3	Tsp		0.3
D70A	34.1	1	PMA	sampled	1
		100			100

PCR: 1 + E2PM21 + RA2 constructs.

1.7. Baic	0.1
12 seeds - 3 No. 1	2
10x	10
10x	10
H ₂ O	3/10
1.00	0.3
10x	1
10x	100

PCR Pym # 42.

Total Volume: 100 L

95°C 5 min
95°C 30 sec
55°C 30 sec } 25x
72°C 1 min
72°C 1 min
U/I

8/3/93

Run Seal of PCR product on
Gel with 1 kb ladder



add equal Volume
to 13% PEG / A.M.C. (500 ul)
then incubate
precipitation at 4°C O/N.

Incubate Seal - LBT Amp/Kan.
for 24 hrs. Inductions
HYPANOL 516 + K₂ + PDIO.
HECm21 + PDEG0 #1
Incubate 37°C w/ aeration 10/N

9/1/95

Spin DNA fragments 10 min
in 70% Ethanol wash

Spin 5 min

Spin 10 min

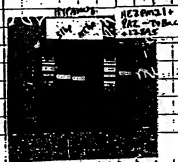
Allow pellet to dry 10 min in RT

Run in on gel 1 kb ladder

Resuspend pellet in 100 ul of

fragments are ready
to digest

Subtract HECm21/H92
H92 + 3 kb fragment
H92 + 10 kb fragment



8/9/95

HIPANX 5162A14 + PDND
HE2PM21 + PQECO #1

Conductions

To 300 ml LBT Amp w/ a ruler
QW Culture fill OD ~ 0.8
Incubator w/ aeration at 37°C
for 2 hrs. fill OD₆₀₀ = 0.4-0.6

Add 100 µg M IPTG to 2 mM ~ 6.5 ml
incubate w/ aeration 4 hrs

at 37°C 15 min
Spin 5K 15 min
Transfer supernatant
resuspended pellet in 10 ml H₂O
HE2PM21 ~ 40 ml
HIPANX ~ 20 ml
Store 4°C over weekend.

9/4/95 Labor Day

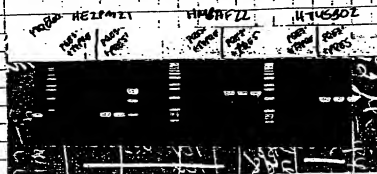
off.

9/5/95

Spin Cultures BK: 20 ml
Transfer supernatant to fresh
tube
Prepare fresh N₂ column
2 ml Buffer B. 2
20 ml 0.1 M N₂ Soy
20 ml 10 mM

9/5/95

Run 10 ul on gel with 11 kb ladder



looks like one of the primers is not working
 - probably #12 (pg 60) T2 pQE60
 made by New England Biolabs
 or Redo.

Digest New HTPANORSON 51bp + 185bp
 pQE60 constructs

PCR Fragment	10
Q10X	5
H ₂ O	34.0
Nico	2.5
BspH	2.5
	50ul

incubate
 37C 0/12.

9/6/95

Precipitate Digests
 Add 150 ul TE = 500 ul 13% PEG/1000
 mix well
 Spin 10 min

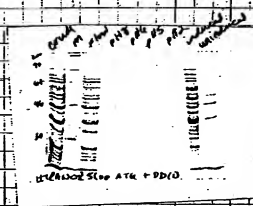
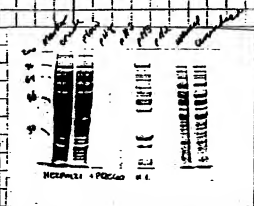
9/16/95

Wash Pellet 70% ethanol 5 min
 Spin 5 min
 Resuspend Supernatant
 Wash Pellet to Dry at RT - 10 min
 Resuspend pellet 1 in 100 μ l TE
 Set up ligation

5bp NIB	4	4	4	4	4	4
185bp NIB	2	2	2	2	2	2
PGE NIB	2	2	2	2	2	2
10X	1	1	1	1	1	1
T4 ligase	1	1	1	1	1	1
H ₂ O	11	11	13	13	15	17

Incubate 110°C O/N.

DE STAIN Gels



Computer Work

9/7/95

Transform E. coli

Thaw 20 min up 5 Cells mice
 100 μ l of Chemically Competent
 cells add 10 μ l (0.1) Negative
 Let sit 10 min 11 min
 Heat 42°C 45 sec
 Place on ice
 Add 400 μ l LB
 Incubate 37°C 1 hr
 plate 100 + 200 μ l onto LB + Amp/Kan
 plate
 Incubate 37°C O/N

- HTPKOR Silp - Ap from P. aeruginosa + Black O/N
 1:200 Dilution in Blocking Buffer

Computer Work -

ORF for HNFAR04
 HTUS02
 HSEK009
 HARS4F22
 HE2UM21

9/8/95

Pick Colonies into 200 μ l LB + Amp/Kan
 in 96 well dish
 185 μ l ATG and 51 μ l ATG
 Incubate 37°C 4 hr @ 190 rpm
 Ready to do PCR to check for
 inserts

140.

9/8/95

PCR to check for HTRAN08.S04
51bp & 185bp ATG

51bp			185bp		
5: 100	2	90x	5: N/A	2	90
3: 100	0.1	90	3: 100	0.1	90
10x	3.2	28x	10x	3.2	28
10x	3.2	28x	10x	3.2	28
10x	21.3	197	10x	21.3	197
10x	0.2	18	10x	0.2	18
10x	2		10x	2	
10x	3.2		10x	3.2	

PCR Program 60

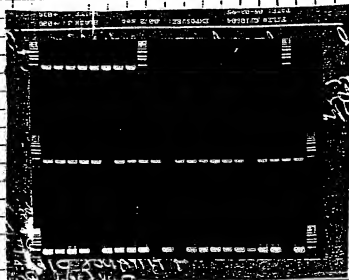
95°C 5 min
 95°C 20 sec
 55°C 20 sec } 30x
 72°C 1 min
 72°C 7 min
 4°C Hold

Run 10.0 on out 1126 2.20

HTRAN08.S04 1126 2.20

HTRAN08.S04 1126 2.20

9/8/95



looks like
transformations
worked very
well

A-H = HTPAN088
51bp ATG + POC

A'-H' = HTPAN089
180bp ATG + POC

Inoculate 3ml LB + Amp/Kan with
10ul of 4hr culture
incubate at RT over weekend
1-2 leng both 51bp + 185bp ATG

FAS Ligand AB. See pg 8/31/95
#11940 & #11941 large bleed.

Pre-ABO Buffer + 1° Ab
Wash 1x PBS 10 min
Add Anti Rabbit Peroxidase (2° Ab)
at 1:2000 in 1x PBS
Incubate at RT 10 min shaking 1 1/2 hrs
Wash filter 1x PBS 10 min
Pierce 50mM Na2HPO4 pH 7 for
5 min
Add Substrate 12.5mM 50mM Na2HPO4 pH 7
25mM A-1000
25mM Phenol
30% H2O2
0.375 ml UBT

90x

180

7

288x

288

1917

18

HTPAN089 1954 ATG + POC 160

9/8/95

Shape cont. color appears to decrease
 brightness by adding dH₂O
 Strip Reaction by adding dH₂O
 Please getters 2x Id H₂O
 Dry on Whatman paper



- Cont.
- 1 HTPANOS 51p 174-175
 - 2 HTPANOS 51p 174/175
 - 3 HUSAT22 174/175-176
 - 4 HTPANOS 51p 174-175
- purified Protein

9/11/95

To the Cultures left at RT mention
 incubated 200mm IPTG to
 2mm - about 4 hours
 incubated 37°C
 spin 10000 x g 5 min
 Resuspend pellet in 75ul H₂O
 Add 15ul 2x detergent
 Buffer 12 ul in 10 to 75ul water
 Run on 12% gel

9/11/95

Run 100V 1/2 hours

STAIN Overnight

#13001-HE2PM21 5' BamHI-Pst
 CCG CCG GGA TCC ACC ATC ATG GCG GCA GCA GTC GTC

PCR New HE2PM21 Pst Construct
 with 5' Bam primer
 #1301

2-Whisk
 4-1000
 2-1000
 4-1000
 in

13101	2
3x dNTP	2
10x dNTP	10
10x PCR	10
H ₂ O	74.8
Tag	0.2
DNA (longer)	1
	100

Total of 1000 µl Run

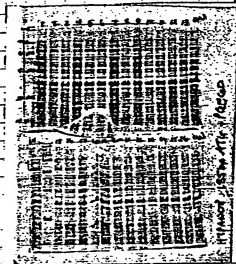
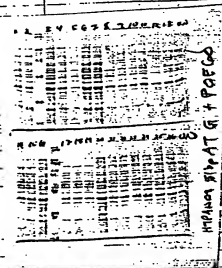
PCR Program TS

95°C	5 min	} 25x
95°C	30 sec	
55°C	30 sec	
72°C	1.5 min	
72°C	7.5 min	
4°C	Hold	

9/12/95

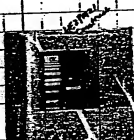
DESTAIN
 HTPACK GUS

9/12/95



looks like H19A108 185 bp ATG induced
 but not 5bp
 Do Baiting peps + Digest
 Geneset 5 min TB + Amp/Km
 with 1-18 g 5bp + 130bp
 incubate 37°C 6 D/A 130bp

H19A108 PA2 Cmatruct
 Run 10 min gel with
 1 kb ladder

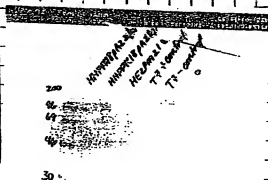


looks good
 PPT with equal volume
 12% PEG / ABC2
 Spin 10 min
 Pour 10 Supernatant
 Wash pellets

146

9/10/95

TNT RESULTS 9/12/95			
INVESTIGATOR	SAMPLE NAME	EXPECTED PRODUCT SIZE KDa	OBSERVED PRODUCT SIZE KDa
MARIO CEPEDA	HTSFH75	30 OR LESS	NO PRODUCT OBSERVED
LAURIE INSCORE	HHFFK18	4.4	48 (major product), 44, 31, 23
LAURIE INSCORE	HHFFK18 PA2 2/25	3.3	NO PRODUCT OBSERVED
LAURIE INSCORE	HHFFK18 PA2 3/18	3.3	NO PRODUCT OBSERVED
ANN KIM	HE2PM21 PA2	4.4	NO PRODUCT OBSERVED
T7 POSITIVE CONTROL	DNASE 02-105	3.3	36 (faint band)
T7 NEGATIVE CONTROL	WATER	NONE	33
T3 POSITIVE CONTROL	HCAC193	3.3	NONE
T3 NEGATIVE CONTROL	WATER	NONE	33



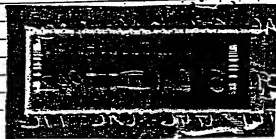
But 1 up PCR of New TNT Controls
using the 0.12uL T7 PCR 100/70
primer (See pg 136 9/5/95).

HE2PM21	HMSAF22	HT45B02	HT45B02
4047	4047	4047	4047
3' size	1.5	1.5	3'
10x dNTP	1.0	1.0	3'
10x PCR	1.0	1.0	3'
H ₂ O	76.2	76.2	3'
Taq	0.3	0.3	3'
DATA	1	1	3'

9/12/95

PEG Pna 106

Run 10ml ongel w/ 1.66kappa



Looks like
only HmsAF22
works well
- try different
program

Tz Parameter is very long
at #4042 - 32ms ...

PEG PPT HmsAF22

9/13/95

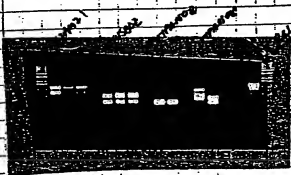
To Rxns from 9/12/95

HE2PM21		HTMS802		516s HTP80008		1080s HTP81008	
4032	0.1	4032	0.1	4032	0.1	4032	0.1
3	1.5	3	1.5		1.5		1.5
10x	1	10x	1		1		1
10x	1	10x	1		1		1
H ₂ O	6.1	H ₂ O	6.1		6.1		6.1
T _{eq}	0.3	T _{eq}	0.3		0.3		0.3
	10		10		10		10

10ml / Tube

Run 10 μ l on Gel w/ 1 kb ladder

8/9/13/95



All PCR Exts
Seems to have
worked.

PEG PPT
- Add Equal Vol
PEG/Wash
- 70% EtOH & Wash
- Repeat 450 μ l TE

Get Cos Cells from Peter Hudson (PH)
from Stock Plate into T25 flask

Boiling media Prep

Spin 2ml culture 5min
Resuspend pellet 750 μ l
3xET + RNase / lysis buffer

Boil 5min

Spin 10min

Remove Pellet

Add 750 μ l PEG/Wash

Mix Well

Spin 15min

Remove Supernatant

Wash Pellet 70% EtOH

1ml

Spin 5min

Remove Supernatant

Resuspend pellet to 150 μ l TE

Run 2 μ l on gel w/ 1 kb ladder

150

9/13/95



Set-up Digests:

DNA	5	36x
10x	3	108
H ₂ O	21.6	777.6
Nco I	0.2	7.2
Bgl II	0.2	7.2
	30 μ l	

Incubate 37°C O/N

9/14/95

Run 2ul of PfuI PfuI
fragments in Gel w/ 1 kb ladder

9/ 9/14/95

Run Balancing Mini prep Digests on
 gel with 1/4 M Laemmli
 10.0 M



Nce / Bal II
 Digest
 All Load
 Lanes only
 Densitometry
 Contract
 Should be done

Run Western of 51 bp induction
 (see pg. 144) 9/12/95

Run 150 V
 Transfer blot 100 V 1 hr

Block 2 hours with Blocking
 Buffer

Probe 10.0 M at 1.200 - Factor
 Add 10.0 M at 1.200 - Factor

Not Bleed # 1199/1200
 Bead 10.0 M / Shaking
 incubate 0.100 M / Shaking

See pg 3 Book # 11
 Lab Note book 405

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